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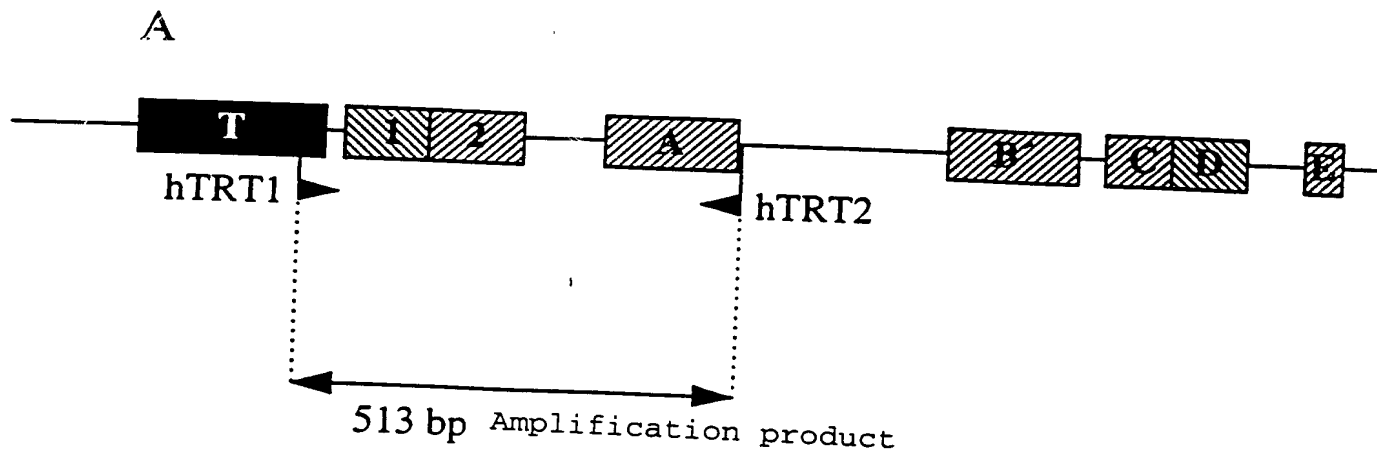
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Fig. 1a

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004030 : 5450960

Fig. 1b



B

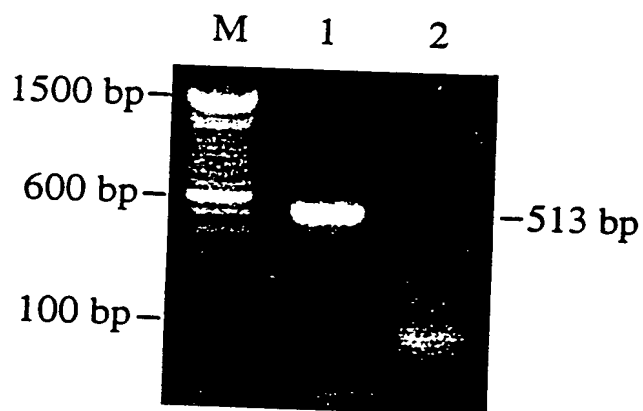


Fig. 2

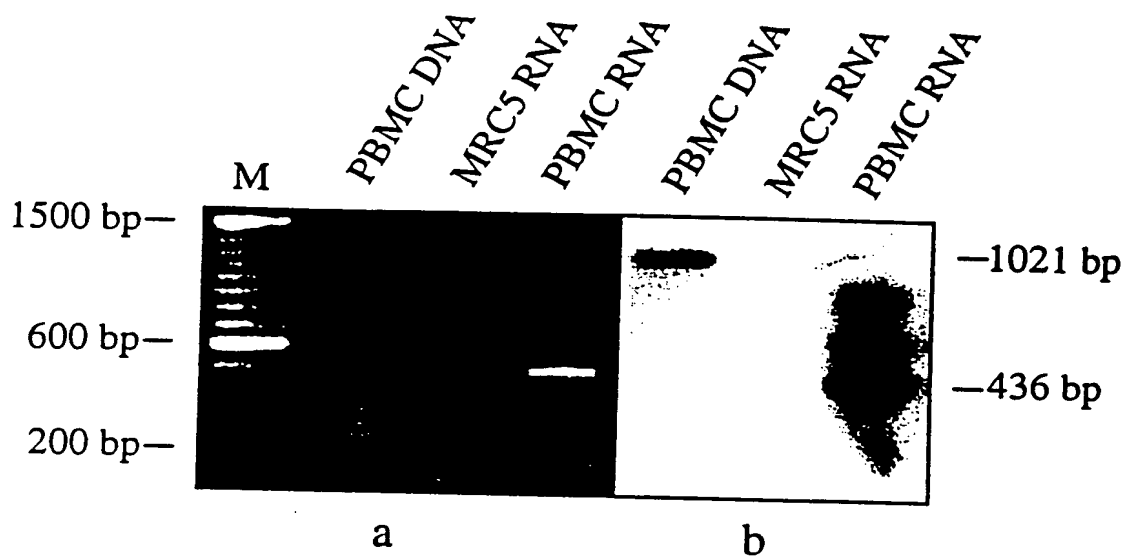


Fig. 3

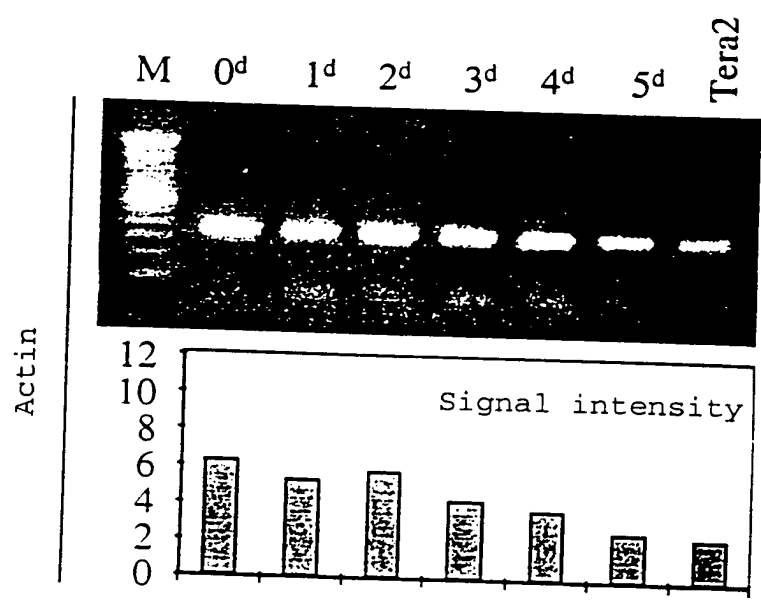
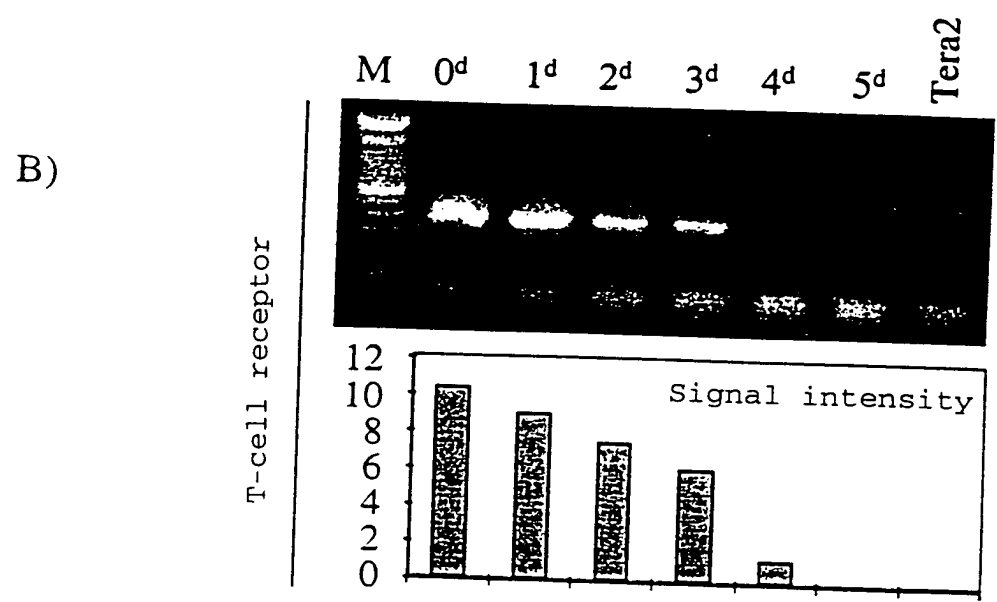
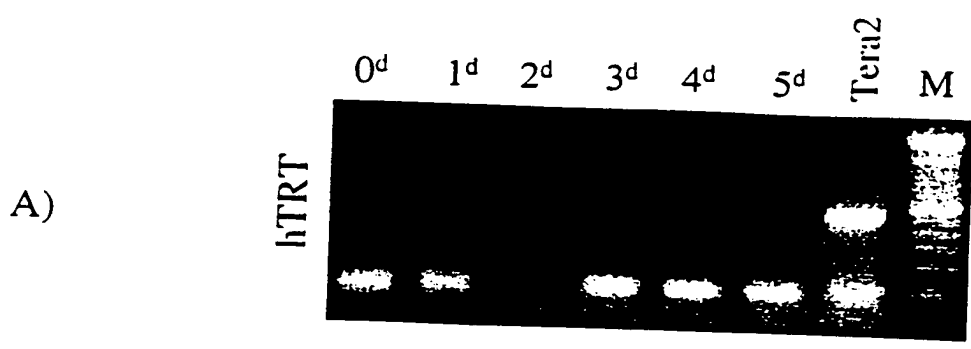


Fig. 4

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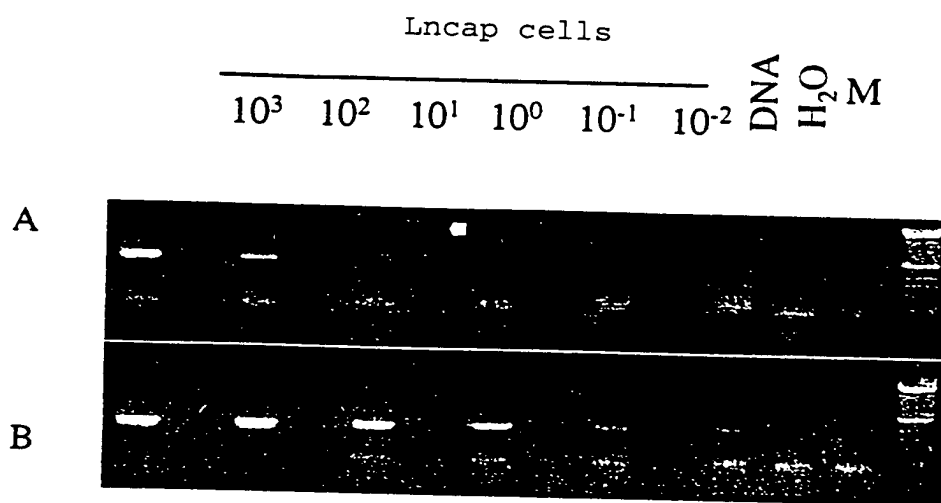


Fig. 5

- 55 -

Catalytic subunit of human telomerase

Name	Sequence (5'-3')
hTERT 1	CTACCGGAAGAGTGTCTGGAGCAAGTTGCAAAGC
hTERT 2	GGCATAACCGACGCACGCAGTACGTGTTCTG
hTERT 3	CGTTCTGGCTCCCACGACGTAGTC

T-cell receptor:

Name	Sequence (5'-3')
TCR 1	GAGGTCGCTGTGTTTGAGCCATCAGAAG
TCR 2	GATCTCATAGAGGATGGTGGCAGACAG

 β -actin:

Name	Sequence (5'-3')
Act 1	GATGATGATATCGCCGCGCTCGTC
Act 2	CTCAAACATGATCTGGGTCATCTTC

 β -globin:

Name	Sequence (5'-3')
Glob 1	ACCCAGAGGTTCTTTGAGTC
Glob 2	TCTGATAGGCAGCCTGCACT

Fig. 6

- 56 -

Whole blood/Ficoll-isolated PBMC

RNA isolation using the 'whole
blood protocol or standard methods
such as phenol/chloroform or silica
gel column

total RNA

Amount of RNA corresponding
to a certain vol. (e.g. 1 ml)
of whole blood

buffer : 100 mM Tris/Cl,
pH 8.3

50 mM KCl
5 mM MgCl₂
1 mM dNTP-mix
2.5 µM random hexamer
4U Rnase-free DNase
40U RNase inhibitor
in a volume of 36 µl

DNase- digest

30' / 37°C, 10' / 75°, 10" / 90°C
→ immediately on ice

sample (18 µl) negative control (18 µl)

+ 50U MuLV RT
+ 40U Rnase inhibitor

+ 4 µl DEPC water

reverse traskriptase reaction

30' / 42°C, 5' / 99°

cDNA

(cDNA)

buffer : 100 mM Tris/Cl,
pH 8.3

50 mM KCl
2 mM MgCl₂
200 µM dNTP-mix
2.5U AmpliTaq DNA
polymerase
300 µM per primer
in a volume of 25 µl

PCR-reaction¹⁾

analysis/quantification

1) PCR-conditions :

15" / 97°C (15" / 97°C, 30" / 70°C, -0.5°C/cycle, 30" / 72°C) x10
(15" / 94°C, 30" / 65°C, -0.5°C/cycle, 30" / 72°C) x20
(15" / 94°C, 30" / 50°C, 30" (ext. 15" / cycle) / 72°C) x10
7' / 72°C

Fig. 7

004080-54910960

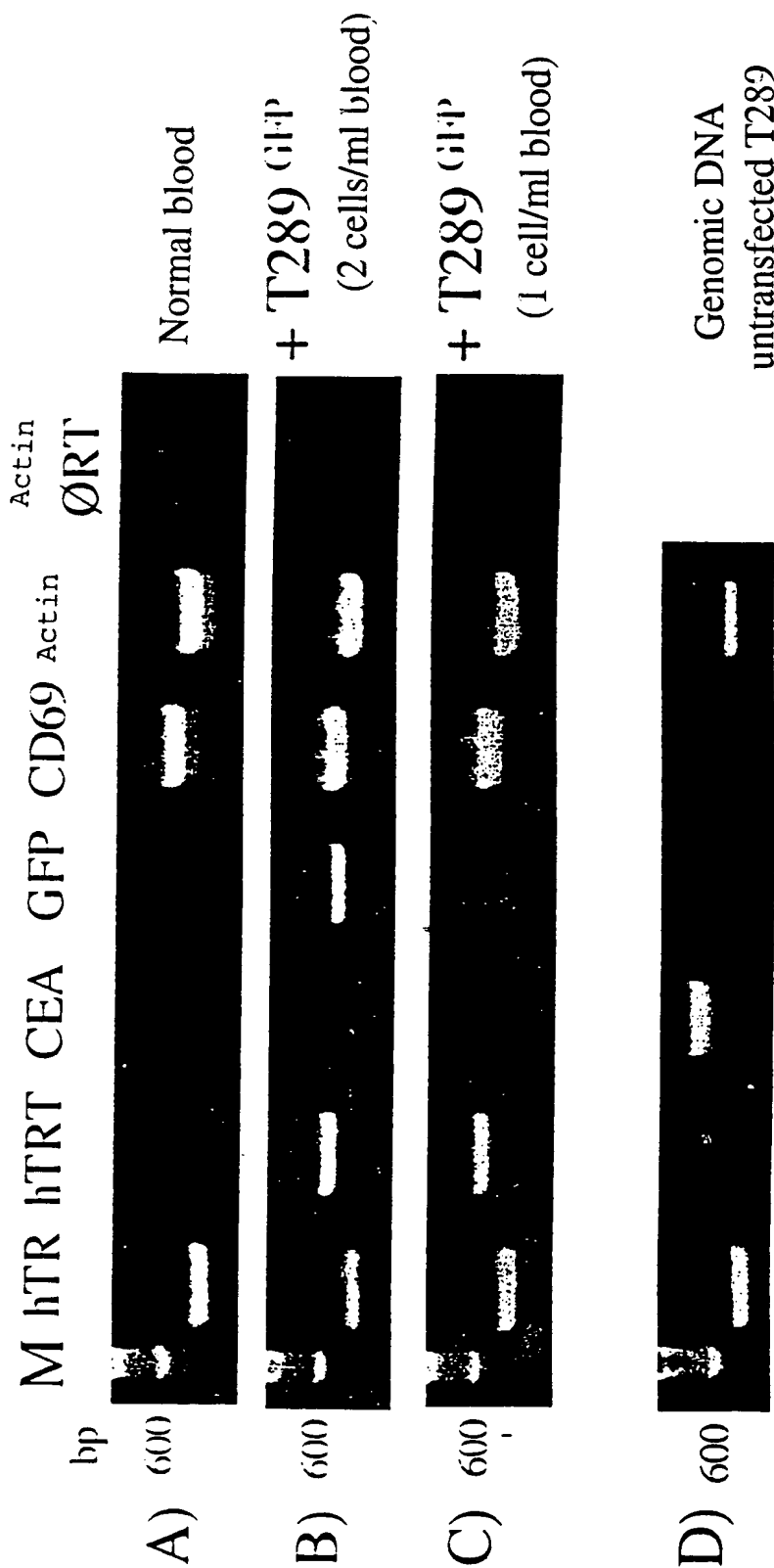


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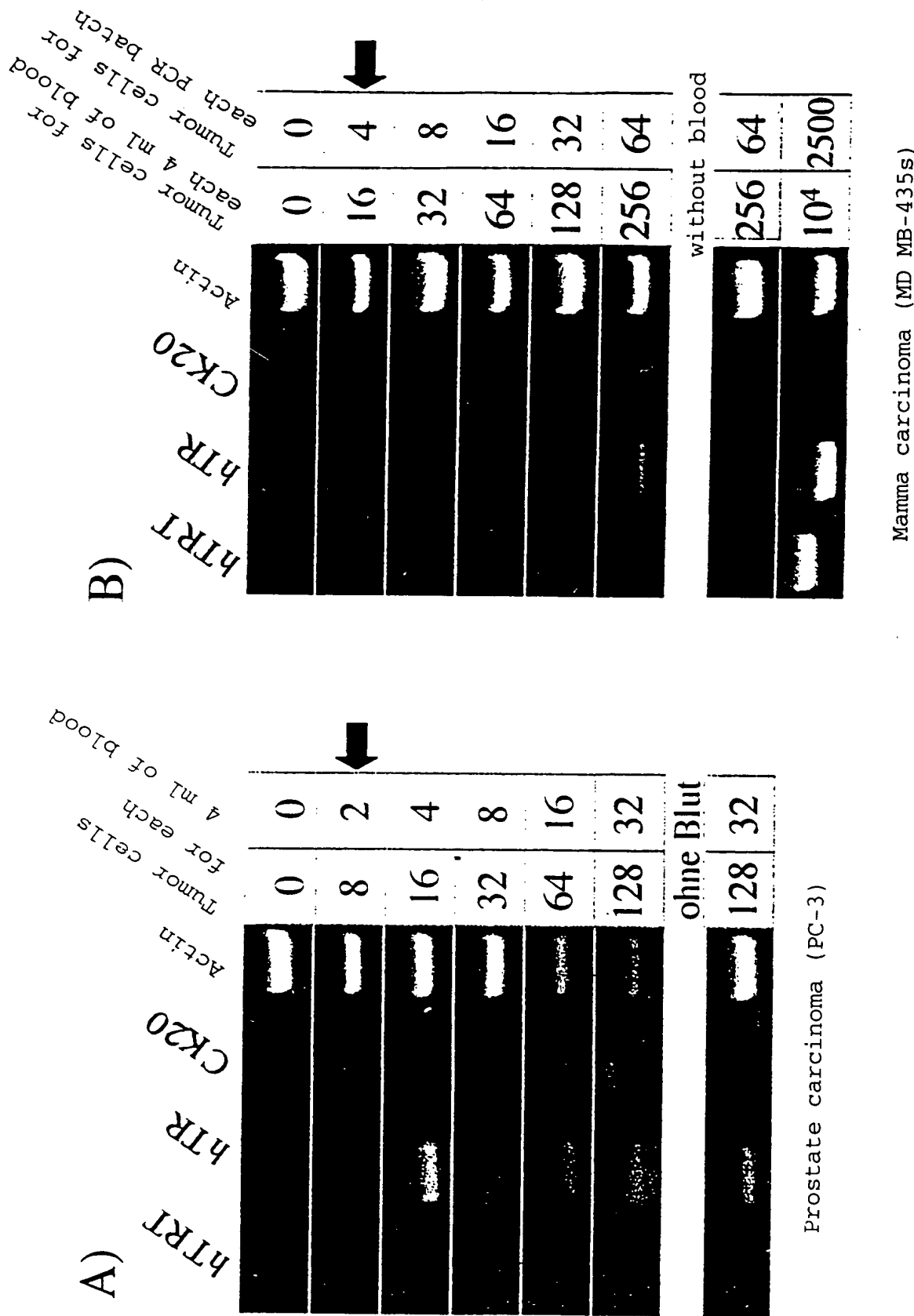


Fig. 9

- 59a -

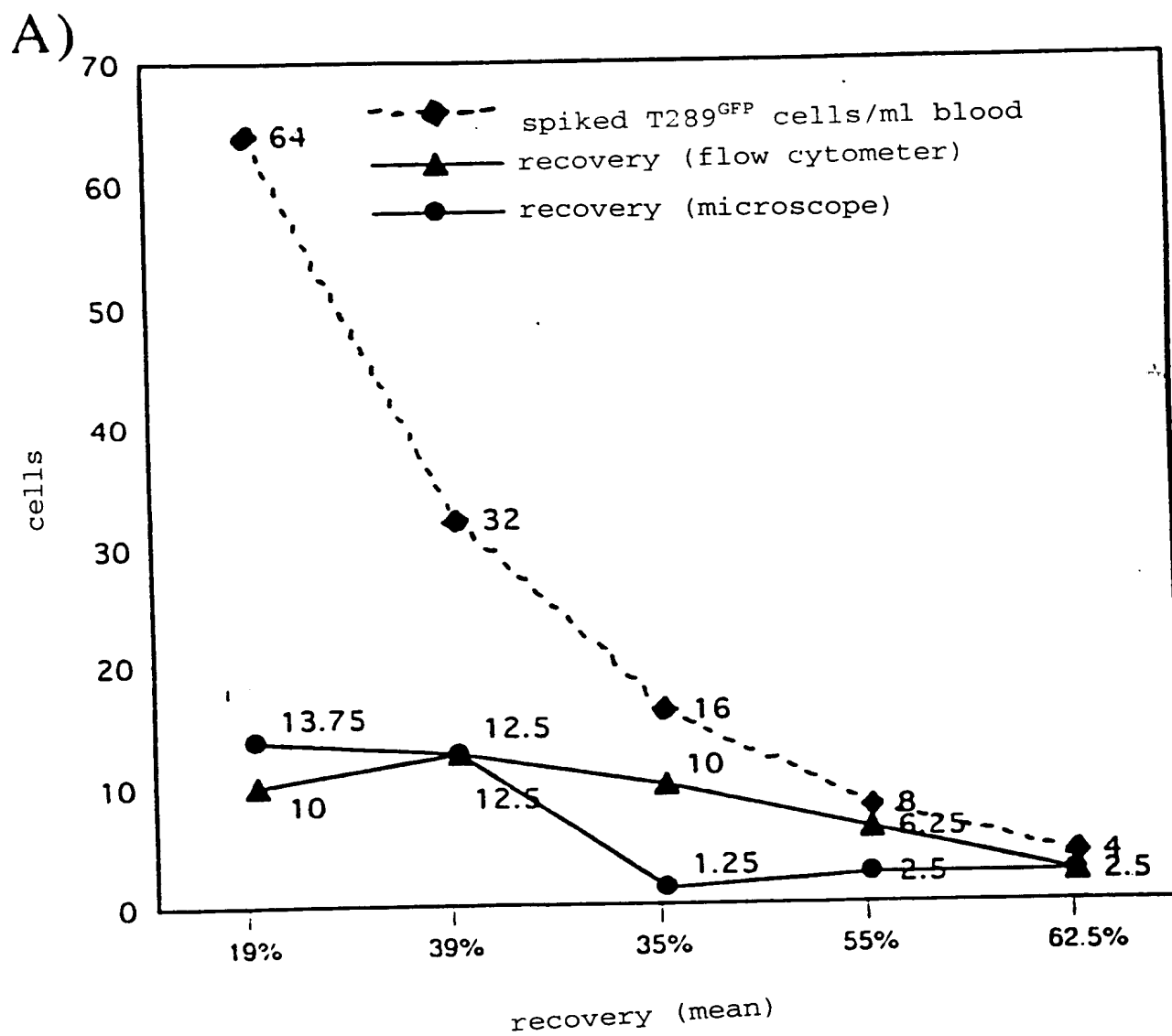


Fig. 10

- 59b -

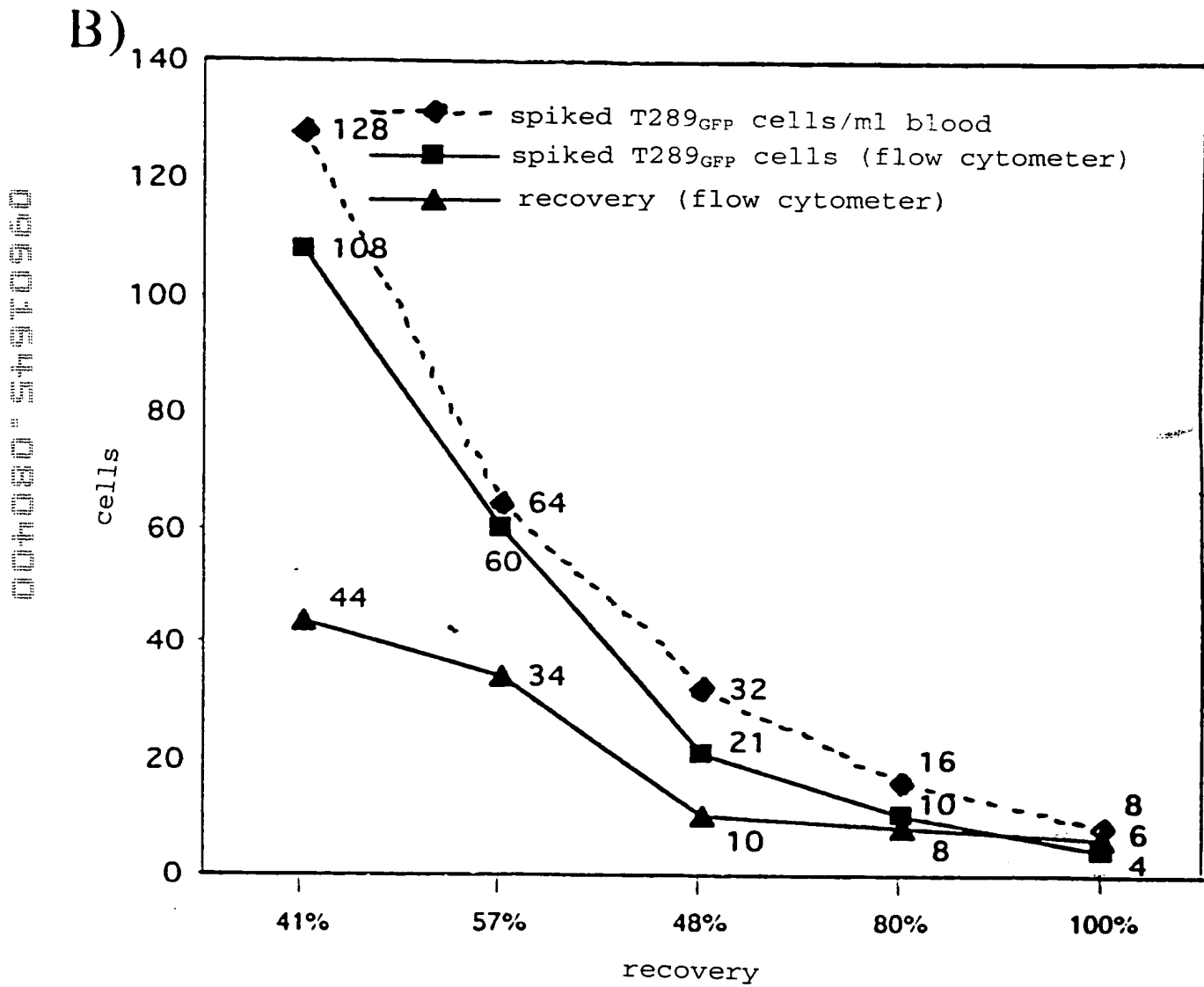


Fig. 10

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